

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information

Center (STIC) detected errors when processing the following computer readable form:

RECEIVED

Application Serial Number: 09/424,70**5**A 0CT **0** 4 2002

Source: 7600

Date Processed by STIC: 9/27/2002 TECH CENTER 1600/2900

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- Hand Carry directly to:
 U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202

U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002



OCT 04 2002

TECH CENTER 1600/2900



Does Not Comply

Corrected Diskette Needec

1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/424,705A

DATE: 09/27/2002 TIME: 13:45:57

Input Set : A:\EP.txt

Output Set: N:\CRF4\09272002\I424705A.raw

4 <110> APPLICANT: LITTLE, MELVYN KIPRIYANOV, SERGEY 6 MOLDENHAUER, GERHARD DEUTSCHES KREBSFORSCHUNGSZEUTRUM 7 ANTIBODY 9 <120> TITLE OF INVENTION: MUTATED OKT3 (ATIBODY 12 <130> FILE REFERENCE: 035280047US00 14 <140> CURRENT APPLICATION NUMBER: 09/424,705A $\Theta \not \mapsto$ 15 <141> CURRENT FILING DATE: 2002-06-02 17 <150> PRIOR APPLICATION NUMBER: PCT/DE98/01409 18 <151> PRIOR FILING DATE: 1998-05-22 20 <160> NUMBER OF SEQ ID NOS: 11 22 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

442 <210> SEQ ID NO: 10 443 <211> LENGTH: (48) 38 shown 444 <212> TYPE: DNA 445 <213> ORGANISM: Homo sapiens 447 <400> SEQUENCE: 10 E--> 448 ccaggggcca gtggatagac aagcttgggt gtcgtttt

450 <210> SEQ ID NO: 11

451 <211> LENGTH: (48) 38 452 <212> TYPE: DNA

453 <213> ORGANISM: Homo sapiens

455 <400> SEQUENCE: 11

E--> 456 cagccggcca tggcgcaggt scagctgcag sagtcwgg

VERIFICATION SUMMARY

DATE: 09/27/2002 TIME: 13:45:59

PATENT APPLICATION: US/09/424,705A

Input Set : A:\EP.txt

Output Set: N:\CRF4\09272002\1424705A.raw

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:448 M:254 E: No. of Bases conflict, LENGTH:Input:48 Counted:38 SEQ:10 L:448 M:252 E: No. of Seq. differs, <211> LENGTH:Input:48 Found:38 SEQ:10 L:456 M:254 E: No. of Bases conflict, LENGTH:Input:48 Counted:38 SEQ:11 L:456 M:252 E: No. of Seq. differs, <211> LENGTH:Input:48 Found:38 SEQ:11